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Seq. ID	Seq. No.	Seq. No.	Seq. No.	Seq. No.	Seq. No.
11	280.0 nt	280.0 nt	280.0 nt	280.0 nt	280.0 nt
211	0.0039	0.0030	.92	1.3013	0.5545
212	0.0471	0.0236	1.18	1.8491	0.5429
213	0.0412	0.0037	1.03	1.8151	0.5505
214	0.0331	0.0002	1.30	1.7300	0.5804
215	0.0076	0.0297	.95	1.8548	0.5451
216	0.0587	0.0075	1.47	1.5785	0.5543
217	0.0519	0.0089	1.30	1.7906	0.5577
218	0.0580	0.0216	.96	1.7857	0.5834
219	0.0533	0.0099	1.31	1.7549	0.5701
220	0.0485	0.0086	1.24	1.7194	0.5838
221	0.0438	0.0092	1.22	1.8709	0.5885
222	0.0486	0.0049	1.22	1.3539	0.7174
223	0.0271	0.0187	.63	1.4430	0.5901
224	0.0216	0.0155	.55	1.4011	0.7136
225	0.0360	0.0249	.95	1.5081	0.5543
226	0.0232	0.0150	.59	1.4827	0.5534
227	0.0357	0.0227	.99	1.5720	0.5681
228	0.0332	0.0018	1.31	1.5595	0.5938
229	0.0357	0.0064	1.39	1.5700	0.5552
230	0.0354	0.0042	1.34	1.5198	0.6174

X

DNA SEQUENCING REQUEST
Microchemistry Department

Requested by Aaron Smith Date _____
 Notebook Reference/assay: 95-1057 P. _____ Project: 95
 Vector: pBc/CMV Host: _____ # of Clones 20
 (Please indicate sizes)
 Names of Sample(s): MB 211-230 (20 samples) (OD₆₀₀ or titer)
 (Please provide OD₆₀₀ and insert size)

MB 211	MC 19209	Human sapiens cDNA 3' end	94% (295/260)	TE
212	19209	Human ABL gene, intron 1b, partial seq.	75% (48/129)	U
213	19210	Lecithin: cholesterol acyl transferase	100% (283/283)	T
214	19211	Human type IV collagenase mRNA	99% (779/281)	JO
215	19212	Human nephropontin mRNA	100% (283/283)	M
216	19213	Ganglioside GM2 activator precursor (human)	96% (236/244)	R
217	19214	Human nephropontin mRNA	99% (281/283)	M
218	19215	Human glycoprotein mRNA	68% (178/258)	M
219				
220	19217	Human nephropontin mRNA	98% (275/280)	M
221				
222	19219	Human lipocortin II mRNA	59% (98/109)	T
223	19220	Human sapiens cDNA clone 53128 3'	97% (88/90)	RI
224	19221	Human osteopontin mRNA	99% (279/281)	JO

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Witnessed & Understood by me, _____

Date _____

Invented by _____

Date _____

Recorded by Aaron T. Smith

Spacing: 0.0



